

tggcgttttt ccataggctc cgccc

SEQUENCE LISTING <110> Enge Duan, Dongshen University of Iowa Research Foundation <120> Adeno-associated virus vectors <130> 875.007US2 <140> US10/054,665 <141> 2002-01-22 <150> US 60/086,166 <151> 1998-05-20 <150> US 09/276,625 <151> 1999-03-25 <160> 14 <170> FastSEQ for Windows Version 4.0 <210> 1 <211> 20 <212> DNA <213> Adeno-associated virus <400> 1 cgggggtcgt tgggcggtca 20 <210> 2 <211> 19 <212> DNA <213> Adeno-associated virus <400> 2 gggcggagcc tatggaaaa 19 <210> 3 <211> 505 <212> DNA <213> Artificial Sequence <220> <223> A synthetic consensus sequence <400> 3 cgggggtcgt tgggcggtca gccaggcggg ccatttaccg taagttatgt aacgactgca 60 ggcatgcaag ctcgaattca tcggtagata agtagcatgg cgggttaatc attaactaca 120 aggaacccct agtgatggag ttggccactc cctctctgcg cgctcgctcg ctcgctgagg 180 ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg ggcggcctca gtgagcgagc 240 gagcgcgcag ctgcgctc gctcgctcac tgaggccgcc cgggcaaagc ccgggcgtcg 300 ggcgaccttt ggtcgcccgg cctcagcgag cgagcgagcg cgcagagagg gagtggccaa 360 ctccatcact aggggttcct tgtagttaat gattaacccg ccatgctact tatctacagc 420 ttgcatgcat gtgagcaaaa ggccagcaaa aggccaggaa ccgtaaaaag gccgcgttgc 480

505

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<213> AAV circular intermediate, clone p81
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                                                                        120
                                                                        180
aggtcgcccg acgcccgggc tttgcccggg cggcctcagt gagcgagcga gcgcgcagag
                                                                        240
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
acttatctac cgatgaattc gagcttgcat gc
                                                                       272
<210> 5
<211> 300
<212> DNA
<213> AAV circular intermediate, clone p79
<400> 5
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tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgcgcgc
                                                                       120
tegetegete actgaggeeg ggegaceaaa ggtegeeega geeegggett tgeeegggeg
                                                                       180
gcctcagtga gcgagcgcgc gcgcagagag ggagtggcca actccatcac taggggttcc
                                                                       240
ttgtagttaa tgattaaccc gccatgctac ttatctaccg atgaattcga gcttgcatgc
                                                                       300
<210> 6
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<212> DNA
<213> AAV circular intermediate, clone p1202
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tgatggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcqaccaa
                                                                       120
aggtcgcccg acgcccgggc tttggtcgcc cggcctcagt gagcgagcga gcgcgcagag
                                                                       180
agggagtggc caactccatc actaggggtt ccttgtagtt aatgattaac ccgccatgct
                                                                       240
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                                                                       272
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<211> 165
<212> DNA
<213> Unknown
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<223> SEQ ID NO:1 of U.S. Patent No. 5,478,745
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ccgggcgacc aaaggtcgcc cgacgcccgg gctttgcccg qqcqqcctca qtqaqcqaqc
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gagcgcgcag agagggagtg gccaactcca tcactagggg ttcct
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<210> 8
<211> 282
<212> DNA
<213> rAAV circular intermediate, clone p79
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tagataagta gcatggcggg ttaatcattg cctacaaaga gcccctagtg atggagtggg
                                                                       120
ccactccctc tettegeega gegegeagag agggagtgge caactccctc actaggggtt
                                                                       180
cctggcagtt aatgattaac ccgccatgct acttatctac agcttgcatg catgtgagca
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aaaggccagc aaaaggccag gaaccgtaaa aaggccgcgt tg
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<211> 345
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taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg aqttgqccac
                                                                        120
tecetetetg egegeteget egetegetea ggeegggega ecaaaggteg eeegaegeee
                                                                       180
gcccggcctc agcgagcgag cgagcgcgca gagagggagt ggccaactcc atcactaggg
                                                                       240
gttccttgta gttaatgatt aacccgccat gctacttatc tacagcttgc atgcatgtga
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gcaaaaggcc agcaaaaggc caggaaccgt aaaaaggccg cgttg
                                                                       345
<210> 10
<211> 276
<212> DNA
<213> rAAV circular intermediate, clone p81
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taagtagcat ggcgggttaa tcattgccta caaagagccc ctagtgatgg agcccggcct
                                                                       120
caccgagcga gcgagcgcgc agagagggag tggccaactc catcactagg ggttccttgt
                                                                       180
agttaatgat taacccgcca tgctacttat ctacagcttg catgcatgtg agcaaaaggc
                                                                       240
cagcaaaagg ccaggaaccg taaaaaggcc qcqttq
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<210> 11
<211> 316
<212> DNA
<213> rAAV circular intermediate, clone p86
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taagtagcat ggcgggttaa tcattaacta caaggaaccc ctagtgatgg agttggccac
                                                                       120
tecetetetg egegeteget egetegetga ggeegeeeeg geeteagega gegagegage
                                                                       180
gcgcagagag ggactggcca actccatcac taggggttcc ttgtagttaa tgattaaccc
                                                                       240
gccatgctac ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga
                                                                       300
accgtaaaaa ggccgc
                                                                       316
<210> 12
<211> 208
<212> DNA
<213> rAAV circular intermediate, clone p87
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taagtagcat ggcgggttac tcattgccta caaagagccc ctagtgatgg aattggaatg
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attcaccctc catgctactt atctacagct tgcatgcatg tgagcaaaaa gccagcaaaa
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ggccaggaac cgtaaaaagg ccgcgttg
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<210> 13
<211> 310
<212> DNA
<213> rAAV circular intermediate, clone p88
<400> 13
gccatttacc gtaagttatg taacgactgc aggcatgcaa gctcgaattc atcggtagat
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aagtagcatg gcgggttaat cattgcctac aaagagcccc tagtgatgga gttggccact
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ccctctctgc gcgctcgctc gctgggcccg gcctcagcga gcgagcgagc gcgcagagag
                                                                       180
ggagtggcca actccatcac taggggttcc ttgtagttaa tgattaaccc gccatgctac
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ttatctacag cttgcatgca tgtgagcaaa aggccagcaa aaggccagga accgtaaaaa
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ggccgcgttg
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<210> 14
<211> 334
<212> DNA
<213> Artificial Sequence
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<223> A synthetic portion of the consensus sequence
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gccactccct ctctgcgcgc tcgctcgctc gctgaggccg ggcgaccaaa ggtcgcccga
                                                                                 120
cgcccgggct ttgcccgggc ggcctcagtg agcgagcgag cgcgcagctg cgcgctcgct
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cgctcactga ggccgcccgg gcaaagcccg ggcgtcgggc gacctttggt cgcccggcct cagcgagcga gcgagcgcc agagagggag tggccaactc catcactagg ggttccttgt
                                                                                 240
                                                                                 300
agttaatgat taacccgcca tgctacttat ctac
                                                                                 334
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